

1631

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/767,764A

DATE: 02/11/2002 TIME: 14:30:30

ENTERED

Input Set : A:\Seq Listing.ST25.txt Output Set: N:\CRF3\02112002\I767764A.raw



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3 <110> APPLICANT: Church, George M.

5 <120> TITLE OF INVENTION: METHOD OF MAKING PROTEIN ARRAYS

7 <130> FILE REFERENCE: 10498-00009

9 <140> CURRENT APPLICATION NUMBER: 09/767,764A

10 <141> CURRENT FILING DATE: 2001-01-23

12 <150> PRIOR APPLICATION NUMBER: 09/522,732

13 <151> PRIOR FILING DATE: 2000-03-10

15 <160> NUMBER OF SEQ ID NOS: 37

17 <170> SOFTWARE: PatentIn version 3.1

19 <210> SEQ ID NO: 1

20 <211> LENGTH: 17

21 <212> TYPE: DNA

22 <213> ORGANISM: Bacteriophage T7

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30 <212> TYPE: DNA

31 <213> ORGANISM: Artificial sequence

33 <220> FEATURE:

34 <223> OTHER INFORMATION: Amplification primer

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45 <220> FEATURE:

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60 <220> FEATURE:

61 <221> NAME/KEY: misc_feature

62 <222> LOCATION: (1)..(1)

63 <223> OTHER INFORMATION: 5' end modified with acrydite

66 <400> SEQUENCE: 4

67 ccactacgcc tccgctttcc tctc

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- 138 <212> TYPE: DNA
- 139 <213> ORGANISM: Artificial sequence
- 141 <220> FEATURE:



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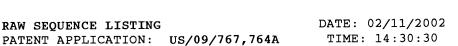
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Input Set : A:\Seq Listing.ST25.txt
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The all numbers has been derected in the Sequence Listing. It view the Sequence Listing to incure a corresponding emplanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

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DATE: 02/11/2002
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Input Set : A:\Seq Listing.ST25.txt
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L:241 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:321 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:339 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:357 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:375 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:393 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:411 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27